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AUTHOR'S CORRECTION

Pathogenomics of the Virulence Plasmids of *Escherichia coli*

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Volume 73, no. 4, p. 750–774, 2009. Page 754: The last two rows of Table 3 should be deleted.

Page 754, column 2: Lines 12–20 should read as follows. "... Like most ETEC strains, H10407 contains multiple plasmids that encode its primary virulence factors. The ensuing analysis of these plasmids will further our understanding of human ETEC plasmid evolution."

Page 754, column 2: Lines 26–30 should read as follows. "... A comparison of pETEC_73 with pCoo (strain C921b-1) reveals that both pCoo and pETEC_73 possess the CS1 operon within a RepI1 backbone region (Fig. 1). ..."

Page 754–755: The first four sentences in the paragraph beginning on p. 754, column 2, line 42, should be deleted. The sentence beginning on p. 755, column 1, line 6, should begin, "Interestingly, the CF-encoding plasmids have ..."

Page 755: Figure 1 should appear as shown on the following page.

Page 767, column 2: Lines 51–57 should read as follows. "... This is particularly evident when E24377A's plasmids are compared to that of strain C921b-1 (pCoo). pCoo forms a cointegrate between the RepFIIA and RepI1 plasmid backbones and contains *eatA* and the CS1 operon (Fig. 1). In 24377A, CS1 ..."

Page 769, Table 4: The entries in the "Replicon(s)" and "Key component(s)" columns corresponding to the pH10407_66 and pH10407_95 rows should read "Not determined."

Page 770, Acknowledgments: Lines 5 and 6 should read, "We give special thanks to Richard Isaacson (University of Minnesota) for critique of this review and to Ian Henderson and Nick Thomson for use of the unpublished sequences of pH10407_66, pH10407_95, and pO42 generated by the Wellcome Trust Sanger Institute."